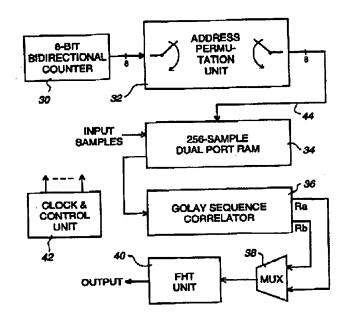
(43) 2001/01/06

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- (51) Int.Cl.<sup>6</sup> H04B 7/216, H04Q 7/36
- (54) SYNCHRONISEUR INITIAL FAISANT APPEL AU CODE DE GOLAY POUR UN CANAL D'ACCES DANS DES SYSTEMES DE COMMUNICATIONS CELLULAIRES
- (54) PREAMBLE USING GOLAY SEQUENCE FOR ACCESS CHANNEL IN CELLULAR COMMUNICATIONS SYSTEMS



(57) A preamble for a reverse access channel (RACH) of a CDMA (code division multiple access) communications system comprises a pair of constituent Golay sequences  $\Lambda$  and B concatenated at least one pair, and preferably a plurality of different pairs, of transformed sequences which are also constituent Golay sequences, each pair of transformed sequences comprising a permutation of elements of the pair of sequences A and B. The permutations can include a reversed order of elements in the sequences, a reversed order of some or all of a set of address bits identifying locations of elements in the sequences, a concatenation of subsets of elements in odd and even locations of the sequences  $\Lambda$  and B, and combinations of these permutations.  $\Lambda$  method for providing the preamble, and related apparatus using a bidirectional counter (30) and an address permutation unit (32), are described.



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## Abstract of the Disclosure

A preamble for a reverse access channel (RACH) of a CDMA (code division multiple access) communications system comprises a pair of constituent Golay sequences A and B concatenated at least one pair, and preferably a plurality of different pairs, of transformed sequences which are also constituent Golay sequences, each pair of transformed sequences comprising a permutation of elements of the pair of sequences A and B. The permutations can include a reversed order of elements in the sequences, a reversed order of some or all of a set of address bits identifying locations of elements in the sequences, a concatenation of subsets of elements in odd and even locations of the sequences A and B, and combinations of these permutations. A method for providing the preamble, and related apparatus using a bidirectional counter (30) and an address permutation unit (32), are described.

## PREAMBLE USING GOLAY SEQUENCE FOR ACCESS CHANNEL IN CELLULAR COMMUNICATIONS SYSTEMS

This invention relates to preambles using a Golay sequence for an access channel, in particular a so-called reverse access channel (RACH), in a cellular communications system.

#### Background of the Invention

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In cellular communications systems, for example in a wireless cellular communications system using CDMA (code division multiple access) techniques for communications between a mobile terminal or station (MS) and a base station (BS), it is well known for the BS to transmit a pilot signal and a broadcast message including a preamble. On being powered up in a cell associated with the BS, a MS uses the pilot signal for synchronization to the BS, and downloads information including the preamble from the broadcast message. Having accordingly determined the timing of the BS, the MS transmits the preamble on the RACH. This is detected by the BS using correlation techniques, so that the BS is informed of the MS, and it can proceed with establishing communications on a traffic channel between the BS and the MS.

In submission TSGR1#3(99)205 to the TSG-RAN Working Group 1 meeting #3, March 22-26, 1999, antitled "New RACH preambles with low auto-correlation sidelobes and reduced detector complexity" (a copy of which is included herein as an Appendix at the end of the description), it is proposed that the preamble, which comprises 4096 code chips providing one of 16 orthogonal signatures of length 16 complex signals, be provided by binary Golay sequences, which have the advantageous property that the sum of their aperiodic auto-correlation functions is zero for all non-zero time shifts.

Consequently, that submission proposes that the preamble be formed from a pair of complementary sequences A and B, which together constitute a Golay sequence and are referred to as constituent Golay sequences, each of 256 chips, repeated in a specific one of 16 signature patterns, so that the overall sequence has a length of 4096 chips, as shown by Table 1 below.

In Table 1, the signature patterns include the sequences A and B in normal and inverted forms, the inverted forms being denoted -A and -B respectively. The 4096 chips of an overall sequence can conveniently be included in one 10 ms time slot, the constituent Golay sequences A and B being individual to a specific cell and/or BS, and the above signature patterns being the same for all cells and base stations.

In contrast to using longer distinctive codes or sequences each of 4096 chips to identify the BS and any of 16 users, for which matched filtering would be required for sequences of 4096 chips, this preamble construction enables the matched filtering to be applied to the much shorter sequences of 256 chips, with a consequent substantial reduction in computational complexity.

1	A	A	В	В	A	-A	- <b>B</b>	В	A	-A	В	-B	A	A	-B	-В
2	Α	A	В	В	A	-A	-B	В	-A	A	<b>-B</b>	В	-A	-A	B	В
3	Α	-A	В	-B	A	A	-B	- <b>B</b>	Α	Α	В	B	A	-A	-B	В
4.	A	-A	В	-B	A	A	-B	-B	-A	-A	-B	-B	-A	A	B	-B
5	Α	A	В	В	-A	A	B	- <b>B</b>	A	-A	В	-B	-A	-A	В	В
6	A	A	В	B	-A	A	В	-B	-A	Α	<b>-B</b>	B	A	A	-B	-B
7	A	-A	В	-B	-A	-A	В	В	A	$\mathbf{A}_{\cdot}$	В	B	-A	A	В	-B
8	Α	-A	В	B	-A	-A	В	·B	-A	-A	- <b>B</b>	-B	A	-A	-B	В
9	A	A	-B	-B	A	-A	В	-B	Α	-A	-B	В	Α	A	В.	В
10	A	A	<b>-B</b>	-B	A	-A	В	-B	-A	A	В	-B	-A	- <b>A</b>	-B	-B
11	Α	-A	-B	В	A	Α	В	В	A	A	-B	-B	A	-A	В	-B
12	Α	-A	-B	В	A	A	В	В	-A	-A	В	В	-A	A	-B	В
13	Α	A	-B	-B	-A	A	-B	B	A	-A	-B	В	-A	-A	-B	- <b>B</b>
14	A	Α	- <b>B</b>	-B	-A	A	-B	В	-A	A	B	-B	A	A	B	B
15	A	-A	- <b>B</b>	. <b>B</b>	-A	-A	-B	<b>-B</b>	A	A	- <b>B</b>	-B	-A	A	-B	В
16	A	-A_	-B	В	-A	<u>-A</u>	-B	-B	-A	-A	В	В	A	-A	B	-B
	Table 1															

It has been found, however, that Golay sequences (and Gold code sequences which have previously been proposed) produce substantial cross correlation peaks for time shifts corresponding to 256 chip boundaries, corresponding for example to a communication distance of about 9 km. Such undesired cross correlation peaks are understood to be due to the repetitive nature of the constituent sequences in the signature patterns discussed above, and present a significant challenge if the delay due to communications distance can correspond to the preamble length or a multiple of the preamble length, and hence for situations where it is desired to use a relatively short preamble and relatively large cell sizes.

Accordingly, aspects of this invention seek to provide an improved preamble, and method for providing such a preamble, for use in an access channel of a communications system, a method of producing an extended sequence from a pair of constituent Golay sequences, and related apparatus.

#### Summary of the Invention

One aspect of this invention provides a method of producing an extended sequence from a pair of constituent Golay sequences A and B, comprising the steps of: transforming the pair of constituent Golay sequences A and B to form at least one pair of transformed sequences which are also constituent Golay sequences; and concatenating at least two of said pairs of constituent Golay sequences to produce the extended sequence.

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The step of transforming the pair of constituent Golay sequences A and B to form each pair of transformed sequences preferably compaises a permutation of elements of the sequences. The permutation can comprise, for example, a reversal of the order of elements in the sequences, a reversal of a set of address bits identifying locations of elements in the sequences, a reversal of some but not all of a set of address bits identifying locations of elements in the sequences, or a concatenation of subsets of elements of the sequences selected by decimation, for example a concatenation of two subsets of elements, in even and odd locations in the sequences, or it can comprise combinations of these permutations.

The invention also provides a method of producing a preamble for use on an access channel of a cellular communications system, wherein the preamble includes an extended sequence produced by the above method. In particular, each of the constituent Golay sequences A and B and each of the transformed sequences can comprise 256 elements, and the preamble can comprise 4096 elements. Advantageously in this case the extended sequence can comprise the constituent Golay sequences A and B and seven different transformations of these sequences A and B.

Another aspect of this invention provides a method of providing a preamble for an access channel of a CDMA (code division multiple access) communications system, comprising the steps of: providing a pair of constituent Golay sequences A and B; permuting elements of the pair of sequences A and B to form at least one pair of transformed sequences which are also constituent Golay sequences; and concatenating said at least one pair of transformed sequences with the pair of constituent Golay sequences A and B for use as an extended sequence in the preamble.

The preamble can be provided by a concatenation of the pair of constituent Golay sequences A and B with a plurality of pairs of said transformed sequences according to different permutations of the elements of the pair of constituent Golay sequences A and B. The step of permuting elements of the pair of sequences A and B can comprise a reversal of the order of elements in the sequences, a reversal of at least some of a set of address bits identifying locations of elements in the sequences, or a concatenation of two subsets of elements of the sequences, the two subsets corresponding respectively to even and odd locations in the sequences.

The invention further provides apparatus for use in carrying out the above methods, comprising a memory for storing elements of each sequence, and address control means for addressing the memory for writing to or reading from the memory consecutively at locations according to the transformed sequences. The address control means can comprise a bidirectional counter for supplying memory addresses incremented in either of two opposite directions, and an address permutation unit for permuting said

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addresses in accordance with the transformed sequences and for supplying the permuted addresses to the memory.

Yet another aspect of the invention provides a preamble for an access channel of a CDMA (code division multiple access) communications system, comprising a pair of constituent Golay sequences A and B and, concatenated with the pair of constituent Golay sequences A and B, at least one pair of transformed sequences which are also constituent Golay sequences, each pair of transformed sequences comprising a permutation of elements of the pair of sequences A and B.

The preamble can comprise a plurality of pairs of said transformed sequences having different respective permutations of the elements of the pair of sequences A and B. In particular, respective pairs of transformed sequences can comprise the pair of sequences A and B with a reversed order of elements in the sequences, the pair of sequences A and B with elements in the sequences permuted in accordance with a reversed order of at least some of a set of address hits identifying locations of elements in the sequences, and elements in odd locations of the pair of sequences A and B concatenated with elements in even locations in the pair of sequences A and B. Brief Description of the Drawings

The invention will be further understood from the following description with reference to the accompanying drawings, in which:

Figs. 1 to 4 are diagrams illustrating various transformations any of which may be carried out in implementations of the invention;

Fig. 5 schematically illustrates in a block diagram stages of a Golay sequence generator known in the art;

Fig. 6 schematically illustrates in a block diagram stages of a Golay sequence correlator known in the art; and

Fig. 7 schematically illustrates in a block diagram a preamble correlator and signature detector according to an embodiment of the invention.

Embodiments of this invention serve to perform various transforms, alone or in combinations, on the constiment Golay sequences that are used to provide a concatenated preamble and signature, in a manner that can eliminate or substantially reduce the undesired cross correlation peaks discussed above. Some of these transforms are referred to as sequence reversal, address bit reversal, partial bit reversal, and resampling or even-odd partitioning, and are discussed in further detail below. The invention is also applicable to other transforms, and combinations of these and other transforms, that provide similar results and that may occur to those of ordinary skill in the art and/or are discussed later below. In each case the transform is applied to the two constituent Golay

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Detailed Description

sequences A and B to result in another two sequences which, it can be shown, are also Golay sequences.

#### Sequence Reversal

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This transform reverses the order of all of the elements of the constituent Golay sequences A and B to produce new constituent Golay sequences  $A^*$  and  $B^*$  respectively. Thus if the sequences A and B are given by:

$$A = [a_1, a_2, ... a_{N-1}, a_N]; B = [b_1, b_2, ... b_{N-1}, b_N]$$

then the sequences A\* and B\* are given by:

$$A^* = [a_N, a_{N-1}, \dots a_2, a_1]; B^* = [b_N, b_{N-1}, \dots b_2, b_1]$$

The preamble and signature can then be formed from both the two constituent Golay sequences A and B and the two reversed sequences  $A^*$  and  $B^*$ , in each case with N=256 chips, as shown by the following Table 2, in which the 16 signature patterns, represented by the normal and inverted (-) forms of the sequences, are the same as in the following Table 2:

-A\* B\* -B\* В -A\* -B\* B\* -B -B A\* 1 **B**\* B\* B -B B\* -A 2 B -B B\* -B\* B -B 3 -B\* B -B\* B В -B\* -B -B\* **B**\* -B -B\* -B A\* -B\* 5 В **B**\* В -B\* B\* В -B\* B 6 В **B**\* -A -A -B\* B\* В 7 -B A\* A A -B -B\* B\* -B\* -B -B B\* -B -B\* 8 A -B B\* -B -B B -B\* 9 A В -B\* -B\* -B -B\* -A B -B\* 10 -B\* B\* B -B\* -B B 11 **B**\* В **B**\* B **B**\* -B B\* 12 -B -B\* **B**\* **B**\* -B В A -B 13 A **B**\* В **B**\* -A B B 14 -B\* B\* B -B -B\* Α B B. 15 -B -B\* B\* B\* -B -A\* -B\* -B 16 -B Table 2

Fig. 1 diagrammatically illustrates this transform for a simple case of 16 elements (N = 16) in a sequence A represented by dots at the top of the illustration, these being reordered as shown by lines into a sequence  $A^*$  represented by dots at the bottom of the illustration.

#### Address Bit Reversal

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This transform reverses address bits, represented as binary numbers from 0 to N-1, which can be used to identify the elements of each of the sequences A and B, to produce new constituent Golay sequences A<sup>r</sup> and B<sup>r</sup> respectively. Thus for example for N = 256, the elements a<sub>1</sub>, a<sub>2</sub>, a<sub>3</sub>, ..., a<sub>254</sub>, a<sub>255</sub>, a<sub>256</sub> of the constituent sequence A can be identified by the binary numbers 00000000, 00000001, 00000010, ..., 11111101, 11111111 respectively, which are reversed to give the binary numbers 000000000, 10000000, 010000000, ..., 10111111, 011111111, 11111111 respectively, corresponding to the elements a<sub>1</sub>, a<sub>129</sub>, a<sub>65</sub>, ..., a<sub>192</sub>, a<sub>128</sub>, a<sub>256</sub> respectively forming the transformed sequence A<sup>r</sup>. the sequence B is transformed to the sequence B<sup>r</sup> in the same manner. This is in effect a random permutation of the positions of the constituent Golay sequences A and B, so that the transformed sequences A<sup>r</sup> and B<sup>r</sup> are also Golay sequences.

As described above in relation to sequence reversal and Table 2, the preamble and signature can be formed from the constituent Golay sequences A and B and the transformed sequences A<sup>r</sup> and B<sup>r</sup>, the latter transformed sequences substituting for the transformed sequences A\* and B\* in Table 2.

Fig. 2 diagrammatically illustrates this transform for a simple case of 16 elements (N = 16). As in the case of Fig. 1, the sequence A is represented by dots at the top of the illustration, and these are reordered as shown by lines into the sequence  $A^r$  represented by dots at the bottom of the illustration.

#### Partial Bit Reversal

This transform is similar to the address bit reversal described above, but reverses only a subset of the address bits, again represented as binary numbers, without making any change in the positions of the other address bits. For example, the most significant one of the address bits may be unchanged for each element of each sequence, with the other binary address bits being reversed in the manner described above, with the result that the transform reorders the elements of the sequence A in two distinct groups to form a transformed sequence A<sup>2r</sup>. The elements of the sequence B are reordered in the same way to produce a transformed sequence B<sup>2r</sup>. A<sup>2r</sup> and B<sup>2r</sup> are again constituent Golay sequences which can be used in place of the sequences A<sup>\*</sup> and B<sup>\*</sup> or A<sup>r</sup> and B<sup>r</sup> as described above. Other partitions between the unchanged and changed address bits can alternatively be made.

Fig. 3 diagrammatically illustrates this transform for a simple case of 16 elements (N = 16). As in Figs. 1 and 2, the sequence A is represented by dots at the top of the illustration, and these are reordered as shown by lines into the sequence  $A^{2r}$  represented by dots at the bottom of the illustration, the reordering in this case being in two groups in

that the most significant bit for each element address is not changed, and the remaining 3 bits for each element address are reversed.

## Resampling or Even-Odd Partitioning

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This transform concatenates two (or more) sub-sequences of the elements of each of the sequences A and B obtained by decimation sampling to produce transformed sequences A\$ and B\$ respectively. The transformed sequences A\$ and B\$ are again constituent Golay sequences which can be used in place of the other transformed sequences as described above. For example, dividing the sequence A into two (even and odd) sub-sequences, its elements al to an are reordered as:

$$A^{\S} = [a_1, a_2, ..., a_{N-1}, a_2, a_4, ..., a_N]$$

and the sequence B is similarly reordered to produce the sequence B\$.

Fig. 4 diagrammatically illustrates this transform for a simple case of 16 elements (N = 16). As in Figs. 1 to 3, the sequence A is represented by dots at the top of the illustration, and these are reordered as shown by lines into the sequence  $A^5$  represented by dots at the bottom of the illustration.

The above are examples of transforms that can be used to permute the elements of the constituent Golay sequences A and B to form new constituent Golay sequences, and it can be appreciated that numerous other transforms can similarly be used. In addition, various combinations of these transforms can be used. For example, the elements of two of such transforms, or of one of such transforms and of the original constituent Golay sequences A and B, can be combined in an exclusive-or (XOR) logic operation to result in further constituent Golay sequences A\* and B\*. For example:

$$\mathbf{A}^{\mathbf{x}} = \mathbf{A} \otimes \mathbf{A}^{\mathbf{r}} = \left[ \mathbf{a}_{1} \otimes \mathbf{a}_{11}, \mathbf{a}_{2} \otimes \mathbf{a}_{12}, \dots \mathbf{a}_{N} \otimes \mathbf{a}_{1N} \right]$$

and similarly for the sequence Bx.

It can be appreciated that although Table 2, and its equivalents using other transformed constituent sequences as described above, provides a reduced repetition of sequences within the overall sequence of  $16 \times 256 \approx 4096$  chips, this repetition can be further reduced by use of different combinations of the transform sequences described above in a hybrid concatenated manner. By way of example, instead of the cycle A, B, A\*, B\* which is repeated four times as shown in Table 2 for each of the 16 signatures (determined by the absence and presence of - signs), a cycle of 16 original, transformed, and repeatedly transformed sequences can be used to provide an effective sequence of 4096 chips.

One example of this is given by the following Table 3:

F1 B1 Ç1 **E**1 G1 HI F G H A1 D1 C D В В A\*T B\*T Ye2 B+2 A<sup>2</sup>r B2r A\*2 B\*2 A\$ B\$ **B**\* Br В **A\*** Table 3

In Table 3, A to H and A1 to H1 represent respective ones of 16 constituent sequences produced as indicated by the second line of the table. For example, the sequence H is constituted by the sequence B after sequence reversal to produce the sequence B\* as described above, this then being subject to transformation by address bit reversal as described above to produce a sequence B\*r. It can be appreciated that the concatenation alternates first the A and B constituent sequences and their respective transforms and then the sequence reversed transformations. Table 3 does not indicate the absence and presence of - signs representing the 16 signatures. Using for simplicity the nomenclature A to H and A1 to H1 of Table 3, the following Table 4 illustrates a possible allocation of 16 Hadamard-Walsh signatures which are arranged to facilitate operation in a high Doppler frequency environment:

1	A	В	С	D	E	F	Ġ	H	A1	<b>B</b> 1	Cl	Dl	E1	Fl	Gl	HI
2	A	-B	C	-D	E	-F	Ġ	-H	A1	-B1	Cl	-DI	El	-F1	G1	-H1
3	A	В	-C	-D	В	F	-G	-H	A1	B1	-C1	-Dl	E1	Fl	-G1	-H1
4	A	-B	-C	Ð	E	-F	<b>-G</b>	H	Al	-B1	-C1	DI	El	-F1	-GI	Hl
5	A	В	C	D	-E	-F	- <b>G</b>	-H	Al	B1	Ci	DΙ	-El	-F1	-G1	-H1
6	A	-B	C	Æ.	-E	F	-Ģ	H	A1	-B1	C1	-D1	-E1	F1	-G1	Hi
7	A	B	-C	-D	-E	-F	G	H	A1	B1	-C)	-D1	-E1	-F1	Gl	Hı
8	A	-B	<b>-</b> C	D	-E	F	G	-H	Al	-B1	-Cl	DI	-BI	FI	Gl	-H1
9	Α	B	C	D	E	F	G	H	-A1	-B1	-C1	-D1	-E1	-Fl	-G1	-H1
10	Α	-B	C	-D	E	-F	G	-H	-A1	Bl	-C1	D1	-B1	F1	-G1	Hl
11	A	В	-Ċ	Œ-	E	F	-G	-H	-Al	-B1	C1	D1	. <b>-E</b> 1	-F1	G1	Hl
12	A	-B	-C	D	E	-F	-G	H	-A1	B1	C1	-D1	- <b>E</b> 1	Fl	G1	-H1
13	A	В	C	D	<b>-E</b>	-F	-G	-H	-A1	-B1	-Ç1	-D1	El	Fl	G1	Hi
14	Α	-B	C	-D	-E	F	-G	H	-A1	B1	-C1	Dl	E1	-F1	Gi	-H1
15	A	В	-C	-D	-B	-F	G	H	-A1	-B1	C1	D1	El	F1	-G1	-H1
16	A	-В	-Ç	D	-E	F	G	-H	-A1	B1	<u>C1</u>	-D1	E1	-Fl	-G1	H1
	Table 4															

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Instead of using transformations of a single pair of constituent Golay sequences A and B as described above to produce longer effective Golay sequences, this can also be done by using another pair of constituent Golay sequences C and D which are not

derivable by transformation from the pair A and B. Thus for example a cycle of four constituent sequences A, B, C, and D can be repeated four times to produce the  $16 \times 256 = 4096$  chips overall sequence length, with the 16 signatures being provided as described above by selective inversion of the individual ones of the constituent sequences. In addition, the transformation techniques described above can be applied to the sequences C and D as well as to the sequences A and B. However, this use of additional sequences C and D results in increased computational complexity and does not appear to produce improved correlation results.

Fig. 5 illustrates two stages 10 and 12 of a known Golay sequence generator which can be used to generate the constituent Golay sequences A and B. Each stage 10 or 12, shown within a dashed line box, has two inputs and two outputs, and the stages are concatenated with the two inputs of the first stage supplied with a binary 1 and the two outputs of the last stage providing the sequences A and B. Although only two stages are shown in Fig. 5, in fact the generator has  $P = \log_2 N$  stages for providing sequences of length N. Thus for sequences of length N = 256 as described above, there are 8 stages of the Golay sequence generator.

The generator stages 10 and 12 all have the same arrangement of a delay unit 14 having an input coupled to one input of the generator stage, a multiplier 15 for multiplying the output of the delay unit 14 by a coefficient W, and cross-coupled adding and subtracting units 16 and 17 for providing respectively the sum and the difference of the other input of the stage and the output of the multiplier 15 to constitute the outputs of the generator stage. The generator stages differ from one another in the delays provided by the delay units 14, these delays being powers of 2 from  $2^0$  to  $2^{p-1}$  and being arranged in any order among the stages, and the weights W which in Fig. 5 are indicated as  $W_1$  and  $W_2$  for the generator stages 10 and 12 respectively. For binary Golay sequences the weights W are binary weights of value  $\pm 1$ .

It is observed that the choices of the weights W and the order of the delay stages of the generator can be selected to provide optimum properties for the resulting constituent Golay sequences A and B.

Fig. 6 illustrates two stages 20 and 22 of a known Golay sequence correlator which is referred to as an efficient Golay correlator or EGC. As in the case of the generator of Fig. 5, the EGC of Fig. 6 has n stages where  $n = \log_2 N$  and N is the sequence length, with each stage having two inputs and two outputs, the stages being concatenated with an input sequence supplied to both inputs of the first stage and correlation outputs Ra and Rb produced at the outputs of the last stage. Each correlator stage includes a delay unit providing a delay from  $2^0$  to  $2^{n-1}$ , a multiplier with a respective weight W of  $\pm 1$ , and cross-coupled adding and subtracting units, in an arrangement

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which is similar to that of the Golay sequence generator except that the delay unit is provided in the other input path.

The use of the EGC of Fig. 6 for the RACH of a cellular communications system are known for example from submission TSGR1#3(99)205 referred to above. In addition, reference is directed to S. Z. Budisin, "Efficient pulse compressor for Golay complementary sequences", Electronics Letters, Vol. 27, No. 3, pp. 219-220, Jan. 1991. As these aspects are known in the art, they are not further described here.

Fig. 7 illustrates a block diagram of a preamble correlator and signature detector according to an embodiment of the invention. Such an arrangement is provided for example in a BS receiver of a cellular communications system, and it can be appreciated that a complementary arrangement, with similar address permutation to perform the transforms as described above, is provided for producing the concatenated preamble and signature.

For constituent sequences of length N = 256, the detector of Fig. 7 comprises an 8-bit bidirectional counter 30, an address permutation unit 32, a 256-sample dual port RAM 34, a Golay sequence correlator 36, a multiplexer 38, a Fast Hadamard Transform (FHT) unit 40, and a clock and control unit 42 which supplies clock and control signals to the other units via clock and control lines which are not shown. Input samples are supplied to an input of the RAM 34 and are stored therein, at addresses provided by the unit 32 via an 8-bit address bus 44 as further described below, in sets of 256 samples corresponding to the Golay sequence length. The samples are read out consecutively from the RAM 34 to the correlator 36, which has the known EGC form indicated above with reference to Fig. 6. The correlator outputs Ra and Rb are supplied via the multiplexer 38 to the FHT unit 38, which serves in known manner to provided a final detector output for example in accordance with the signatures of Table 4 above.

These signatures are of length 16, so that over the duration of the preamble of 4096 chips 16 sets of 256-chip samples are processed in the units 36, 38, and 40 in a substantially known manner. The units 30 and 32, operating under the control of the clock and control unit 42, provide a corresponding synchronized cycle of 16 address permutations for these sets, in accordance with the transforms described above and for example in the manner represented in Tables 3 and 4.

To this end, the address permutation unit 32 is arranged to provide a cyclic selection from among different sets of connections between its inputs and its outputs, as is represented diagrammatically in Fig. 7 by selector switches at the (8-bit wide) input and output of the unit 32. Within the unit 32, these connections can be in the form of direct connections between the inputs and outputs corresponding to the illustrations of Figs. 1 to 4 and their variants as described above, for example as given by Table 3. The sequence reversal transforms (indicated by the \* in Table 3) are conveniently effected by reversing

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the count direction of the counter 30, so that samples are stored in the RAM 34 at a reversed sequence of addresses from that from which they are read out to the correlator. For the other transforms (e.g. A<sup>r</sup>, A<sup>\$</sup>, and A<sup>2r</sup> in relation to the sequence A) represented in Table 3, respective sets of direct connections are selected cyclically by the unit 32 in accordance with Figs. 2, 4, and 3 respectively, in comparison to straight-through (i.e. not permuted) connections for the sequence A.

Accordingly, the various transforms represented in Table 3, or other transforms that may be used, can be easily provided by an arrangement such as that of the bidirectional counter 30 and address permutation unit 32.

It can be appreciated that although Fig. 7 and the above description relate to permuted storage in, and consecutive read-out from, the RAM to provide decoding of the transformations described above, these operations could be interchanged with equivalent effect. In addition, it can be appreciated that substantially the same operations can be provided in substantially equivalent manner for effecting the transformations in generating the transformed sequences. Thus for example the correlator 36 of Fig. 7 could be replaced by a Golay sequence generator, having the form of Fig. 5, whose output is supplied to the RAM, an output of the RAM providing the generated and transformed sequences, with a similar counter and address permutation to that shown in Fig. 7.

Although particular embodiments of the invention and various alternatives have been described in detail, it should be appreciated that numerous modifications, variations, and adaptations may be made without departing from the scope of the invention as defined in the claims.

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#### APPENDIX

TSG-RAN Working Group 1 meeting #3 Nynāshamn, Sweden March 22-26, 1999 TSGR1#3(99)205

Agenda Rem:

Source:

Ericsson

Title:

New RACH preambles with low auto-correlation sidelobes

and reduced detector complexity

Document for:

#### 1 Introduction

The prescrible part of the random access burst signal former proposed for UTRA/FDD has the length of 4096 code okips [1]. The prescrible consists of a signature of length 16 complex symbols, which are spread by a common, 256 chip long Orthogonal Gold sequence called prescrible spreading code. In total there are 16 different signatures, obtained from the orthogonal set of binary Orthogonal Gold sequences of length 16, by multiplying each binary code with the constant complex number  $C=(1+\beta)\sqrt{2}$ , where  $\mu=\sqrt{-1}$ .

The UE transmissions of the readom access bursts can start at a number of well-defined time offsets (access slots), which are synchronised to the frame sync of the primary CCPCE. The primary CCPCE frame sync is cattracted after the cell search procedure in the UE. Therefore the random access presurbles are mostived at the base station at the beginning of each access slot interval with the time uncertainty equal to the round-trip propagation delay.

The current random access preamble construction allows simplified realisation of the bank of consistors required in the base studies random access receiver if this time uncertainty is smaller than 255 chips. However, the speciatic auto-correlation sidelohes of such codes are rather high, which means that the RACE preamble might be detected at wrong time positions. In other words, the preamble detection probability at cornect time positions is detectorized for moderate to high algoral-to-color ratios. Therefore it is desirable to find another random access preamble construction, which would also produce an orthogonal ratio of preamble codes with much lower aperiodic auto-correlation sidelohes, facilitating an afficient matched filter implementation.

## 2 Golay complementary sequences

The new RACH prescribles are based on the application of binary sequences from the Golsy complementary pairs. The major property of the binary sequences from the Golsy complementary pair is that the sum of their specialis and—correlation fractions equals zero for all non-zero time thirts. The Golsy sequences can be constructed for any length  $L=2^N$ , where N is any positive imager, and also for lengths 10 and 26, or for any combination of those three lengths. Besides the complementary property, such sequences axhibit some additional properties which make them attractive as synchronisation codes: they have low specialise auto-contilation sidelobes, and there is a large number of them for a given code length.

If the sequences are of length  $L=2^N$ , there is a general method for the construction of polyphase complementary pairs of sequences, where the Golsy complementary sequences are just a special, binary case. That general construction is defined by the following recursive relation [2].

 $a_{\theta}(k) = \delta(k)$   $b_{\theta}(k) = \delta(k)$   $a_{\theta}(k) = a_{\theta + 1}(k) + W_{\theta + 1}(k \cdot D_{\theta})$  $b_{\theta}(k) = a_{\theta + 1}(k) - W_{\theta} \cdot b_{\theta + 1}(k \cdot D_{\theta})$ ,

$$k=0, 1, 2, ..., 2^{n}-1,$$
  
 $n=1, 2, ..., N,$   
 $D_{n}=2^{N_{n}},$ 

-Arre

 $a_*(k)$  and  $b_*(k)$  are two complementary acqueaces of length  $2^N$ , B(k) is the Kronecker delta function, it is an integer representing the time scale, a is the integer representing the time scale,  $D_a$  is a delay,  $D_a$  is a delay,  $P_a$ , n=1,2,...,N, it any permutation of numbers  $\{0,1,2,...,N-1\}$ ,  $W_a$  is an orbitrary complex number of unit magnitude.

If We has values +1 and -1, the binary (Golsy) complementary sequences are obtained [3].

An efficient metched filter disophy countrionding to the complementary sequences  $a_k(k)$  and  $b_k(k)$  defined by (1) is given in Figure 1. This filter performs the councilation of lapter signal r(k) absolutences by with the two complementary sequences  $a_k(k)$  and  $b_k(k)$ . The two matched filter outputs produce the two counsponding spatiodic cross-countiation functions  $R_{n}(k)$  and  $R_{n}(k)$ . Such a digital filter will be called the Efficient Golay Councilator (EGC), although it is notually the filter matched also to any polyphase complementary pair defined by (1). The matched filter has complex conjugated coefficients  $W_{n}$ , denoted as  $W_{n}$ .

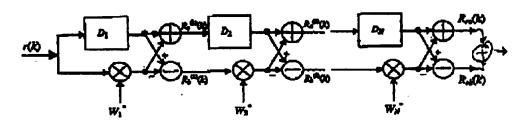


Figure 1: Efficient Golay Correlator (EGC).

The boxes in Figure 1 represent the corresponding delay lines with  $D_s$  memory elements. The number of such discussions in the EUC is equal to  $log_s(L)$ , while in the straightforward matched filter implementation it would be L. The number of additions in the BGC is  $2 \cdot log_s(L)$ , while in the straightforward matched filter implementation it would be L-1. The number of memory elements required for the BGC is L-1 ( $mD_1+D_2+...+D_d$ ), the same as for the straightforward implementation of a single matched filter corresponding to one of the complementary sequences.

## 3 Efficient Golay correlator with reduced memory

In the case when the expected delays  $\tau$  of input signal are limited to be  $tr \in T_{max}$  chips. It is possible to derive another lifticient Goiny correlator with reduced memory. The EGC with reduced memory is based on the representation of a Goiny sequence of length  $L = L^* = L^* - L^* = L^*$  in the so-called "factored" form, i.e. as a function of t = 0 shouter constituent complementary sequences A(k) and B(k) of length  $T_{max}$ . This relation is a simple consequence of the general recursive construction (1), which can actually start from any complementary pair of sequences. Namely, if the initial vectors  $a_0(k)$  and  $b_0(k)$  are taken to be

$$a_i(k) = A(k),$$
  
 $b_i(k) = B(k), \quad k=0, 1, 2, ..., T_{max}-1,$  (2)

where A(k) and B(k) are the two arbitrary complementary sequences of length  $T_{min}$  the resulting pair of complementary sequences of length  $L_m(k) = J \cdot T_{min}$  is generated after J iterations. Note that all the delays  $D_n$  in (1) should be multiplied by the length of contributes sequences ( $T_{min}$ ).

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For comple, if the constituent sequences are of length  $T_{max}=2.56$ , the permutation vector  $P_a$  and the weighting vectors  $W_a$  are of length 16 if the resulting complementary pair should be of length 4096. The resulting Colay sequences consists of 8 times repeated sequences A(k) and B(k), which are multiplexed according to some equivalent binary "instricating" function  $I_0(k)$  depending on the permutation vector  $P_a$ . The orthogonal set of 16 Golay sequences of length 4096, having the constituent sequences A(k) and B(k) of length 256 (and a common interdeaving function), one be obtained by choosing a single permutation vector of length 16, along with 8 appropriately choose weighting vectors.

The additional 16 orthogonal Golay sequences of length 4096 having the same constituent sequences A(k) and B(k), but interleaved according to another interleaving function  $I_1(k)$ , can be obtained by taking

$$d_{\ell}(k) = \beta(k),$$
  
 $d_{\ell}(k) = A(k), k=0, 1, 2, ..., T_{max}-1,$  (5)

and using the same permutation and weighting vectors as for the first set of 16 sequences. Therefore in total there are 32 orthogonal Golay sequences of laugth 4096 having the bommon constituent sequences of laugth 256. In principle it is possible to generate 2/ such orthogonal Golay sequences, where Jal/Inna

The set of 16 orthogonal Golsy sequences of length 4096 which is obtained according to the above algorishm and which can be used in UTRA/FOD is given in Table 1, as a function of two shorter, constituent complementary sequences A(k) and B(k) of length 256. The additional set of 16 orthogonal Golsy sequences of length 4096 can be obtained by replacing A(k) with B(k) and B(k) with A(k) in Table 1. As A(k) and B(k) are orthogonal, the additional set is orthogonal to the first one.

Table 1: First 16 exthogonal Golay sequences of length 4896.

			WE 4-													<b>AU</b>
k	0.35											- 1		- 1		
5(0,4)		A	₽.	В	4	4		В	_4	-4	_21	-8		-41	-	_
SLIS		7	3		A	~4			4		-81		-4	-41	_2	- 2
512.40	1 A	-4	B	B	A	A	-		A	Λ		_8.1	-41	-4	-2	
30.5	┲╬╢	-		-35	4	<b>A</b> .	-7	4	-4	*			-4	-41		-2
24.6		A	-	3	4	A	8	-8		-1	В		-4-	-1	_Ł	-
SCSJO	1-2		8	A-	1	A	B	-38	<b>.</b>	A		7			<u>-0</u>	-8
5(6,8)	1-9-	-	-	- 32	- 4	- 3	8	B	A	A			1	A	3	-3
	1-7	-	- 7	-		-4		3	-4	1	4	-		4	-	
3(7.4)	+-7-	7	-	-3	Ā	4	В	3	A	4	4	4	A	A	8	- 8
3(8,4)	1 4	-	-						4	A	B	4	-A	4	7	-8
2027	1 2	<del>- ?-</del>		72	<del></del>	-	<b>-</b>		A		-8	-3	A	*	B	-8
201019	+ +	-~		-	-	A		-	1	-4	4	-	A	A.	-	1 8
20170	1 7	<del></del>	3	- 5	4	-	1	10	A	4	-3	B	-4		-8	-3
1(13.4)	<b>↓</b>	<u> </u>	_	_	- 74	1-2	13	7	1 3	1 7	9	3	A	$\lambda$		
\$03.10	1.4	ĻĄ	-8	1-3		<del>Ĭ~</del> Ŷ		3	17	7	1	-	3	A	-8	_
2147	1.4	1-4		- B	14	1-2	1-2	1		13	<del>- 1</del>		7	-	1 8	3
3035.83	1 4	-4	-8	4_		- 40										

The BACH prescribe correlator with reduced memory, corresponding to the above set of 32 orthogonal prescribes is shown in Figure 2. The number of memory elements per received signature in this scheme is the same as for the RACH prescribe correlator described in UMTS XX.07. However, the total number of adders and multipliers is significantly reduced due to the one of BGC instead of prescribe apreaching code matched filter.

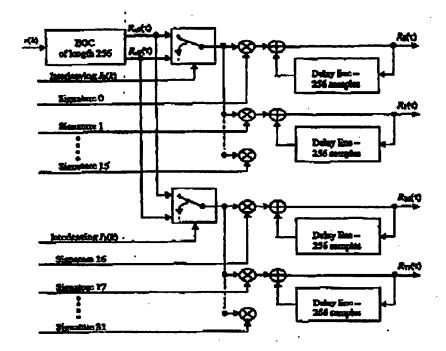


Figure 2: The bank of RACH pressable correlators with reduced memory, matched to 32 orthogonal Galay sequences of length 4096.

The inveriencing function  $I_0(k)$  is common for the 16 orthogonal prombles, while the intergencing function  $I_1(k)$  is common for the other 16 orthogonal premulties. From Table 1 it can easily be seen that

$$I_0(k) = \{0, 0, 1, 1, 0, 0, 1, 1, 0, 0, 1, 1, 0, 0, 1, 1, 1, 0, 0, 1, 1, 0,$$

Each preamble has a unique "rigusture" requence, which can also be easily derived from Table 1. For example,

The set of 256 cell-specific pairs of constituent Golay sequences A(k) and B(k) (corresponding to the set of 256 cell specific prescrible specifing codes) is defined by (1), where the pergramation vector  $P_n$  is common for all pairs and in given by

$$P_{n} = \{0, 2, 1, 5, 5, 4, 7, 3\},\tag{4}$$

while the corresponding 256 weighting vectors  $W(\nu, n)$ ,  $\nu = 0$ , 1, ..., 255, are defined as the 8-bit binary representations of integers  $\{0, 1, 2, ..., 255\}$ , i.e.

$$W(\nu,n) = (-1)^{R_{\alpha}(\nu)}, \quad \nu = 0, 1, ..., 255, \quad n = 1, 2, 3, ..., 8,$$
 (5a)

where B\_(x) is the seth bit in the 8-bits long binary representation of some positive integer x, i.e.

$$x = \sum_{p=0}^{8} B_{+}(x) \cdot 2^{p-1}. \tag{5b}$$

Note that all 256 constituent pales can be detected by using the cases correlator shown in Figure 1, by adopting only the weighting coefficients  $W_n$ .

## 4 'Implementation complexity

The implementation complexity of the heak of RACH promible correlators is significantly reduced due to the use of RGC instead of preschip spreading code metabol filter. Assuming that the number of new orthogonal preembles based on Golsy complementary sequences (GCS) used is the same (16) as in the case of the current preembles based on connected orthogonal Gold sequences (GGS), the implementation complexity of the corresponding banks of correlators can be compared in the following way:

- a) The number of address is 32 (=16+16) for the GCS, compared to 271 (=255+16) address for the GGS.
- b) The number of multipliers is 24 (#8+16) for the GCS, compared to 272 (#256+16) multipliers for the OGS.
- c) There is a multiplexes (swinth) for GCS while there is no multiplexes for OGS.
- d) The lengths of dolay lines are the same in both cases.

## 5 Aperiodic autocorrelation properties

Besides the improved implementation officiency, the new preembles based on Golay sequences offic much better performances in term of the mechanic absolute speciation autoconvolution sidelobes (MAS), when compared with the current proemble codes based on concatenated orthogonal Gold sequences.

As a first countple, the aperiodic auto-correlation function for one of the concatenated orthogonal Gold presumbles generated with the old scheme for presumble apreading code n=1 is shown in Figure 3 in the annex. This should be compared with the new construction using Golay sequences with constituent requences A and B defined by (1), (4) and, as an example,  $W_{n}=\{1, -1, 1, -1, 1, -1, 1\}$ . The speciation suito-correlation function of this new code is shown in Figure 4 in the same A. As can be seen the Golay sequences have much better same-correlation properties.

The MAS for all the promibles based on the above pressible spending code are listed in Table 2. The benefits of the Golsy sequences in terms of reduced MAS is closs.

Table 2: MAS for preambles corresponding to one particular preemble spreading code.

Golay saguence	)eş	Conceinness Orthogonal Gold seguences					
Number of	MA8	Number of occurrences	XAS				
4	161	1	1024				
4	jái	4	1296				
8	133	7	1536				
•		2	1793				
-	•	2	2048				

The random access preambles are not completely asynchronous to the base station receiver because the UE has the basic information about base station timing, but with an uncertainty introduced by the round-stip propagation delay between the base station and UE. The convent assumption in UTRA/FDD is that the round-trip delay is at most 255 chips to be able to the proposed simplified receiver structure, so the speciodic auto-correlation function of rendom access presumbles is accessly of most interest only in the region +/- 255 chips around the main lobe. The maximum absolute values of speciodic surccorrelation sidelobes in the region +/- 255 chips around the main lobe are shown in Table 3 for the previously described Golsy and concatenated Outhogonal Gold sequences of length 4096.

Table 3: MAS in the +/- 255 chips region for preambles corresponding to one particular preamble spreading code.

Golay soquear	- ing	Constitutional Orthogonal Gold sequences					
Number of occurrences	MAS	Number of occurrences	MAS				
16	31	1	731				
	-	3	73/7				
-		3	743				
	-	3	755				
	-	6	761				
		1	167				

From Table 3 it can be nodeed that Golsy requestes have about 25 times lower auto-correlation sidelobes than the concurrenced Orthogonal Gold sequences, in the region +/- 255 other around the main lobe.

It is clear that for the particular codes evaluate above, the Golsy sequences are superior. Finally, the maximum absolute values of speciotic anto-correlation sidelebas in the region +4-255 chips around the main lobe are evaluated for all presumbles. Both the Golsy based 256 pairs of constituent sequences A and B defined by (4) and (5) for all 32 orthogonal presumbles of length 4096 corresponding to each such pair of constituent sequences, and the current presumbles based on concataunted Orthogonal Gold sequences have been investigated. The results are shown in Table 4.

Table 4: MAS in the +/- 255 chips region for all presmbles.

Golsy sequent	<b>A4</b>	Concernment Orthogonal Gold				
Number of	EAM	MAS				
64	27	MAS values are plotted in				
128	29	Figure 5 in the amer.				
1280	31					
1024	33	Averaga MAS is 669, largast				
1600	345	MAS is 1080, smallest MAS is				
1280	37	286.				
832	39					
512	41	95% of MAS values are above				
576	43	<b>560</b> .				
256	45					
320	47					
192	49					
- 64	51					

Table 4 shows that all \$192 possible Golay preambles of length 4096, have corremely low maximum auto-correlation sidelobes. The average MAS is 37, and 65% of the MAS values are between 27 and 37. A simple, but rather fair, comparison between the two different preamble designs can be done by comparing the average MAS. The old concatenated orthogonal Gold preambles have an average MAS 18 times (669/37) higher than the Golay based preambles.

#### Conclusion 6

A nate set of RACH promphles is proposed for inclusion in UTRA/FDD. The benefits of the premphles codes, based on Golsy complementary soquences, are:

The new preambles office significently more efficient preamble detector bandware implementation, measured in terms on the number of multiplions and adders required.

The number of available preembles is doubled, to 5192.

All 8192 of the new prescribles have good sute-correlation properties, while the span for the old preembles is quite large and many of those codes exhibit very bad correlation properties.

The new preembles have about 18 times lower speciadio suite-correlation sidelobes than the present RACH preembles, offering potentially better Eb/No performance.

### References

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M.J.E.Golsy, "Complementary Series", IRE Trans. on Information Theory, Vol.IT-7, pp.82-87, April 1961.

## Annex - Figures

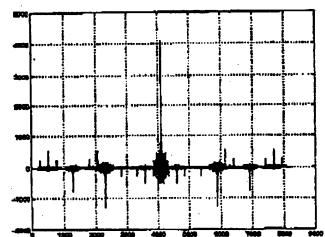


Figure 3: Aperiodic auto-correlation function for one of the present RACH presembles (signature + presmble spreading cods).

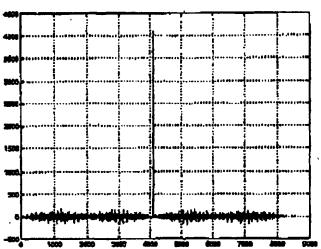


Figure 4: Apariodic auto-correlation function for one of the new RACH preembles (Going complementary sequence from Table 1)

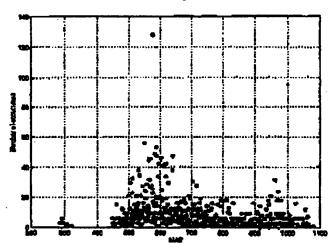


Figure 5: Distribution of MAS values for the old orthogonal Gold hased presembles.

#### WHAT IS CLAIMED IS:

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1. A method of producing an extended sequence from a pair of constituent Golay sequences A and B, comprising the steps of:

transforming the pair of constituent Golay sequences A and B to form at least one pair of transformed sequences which are also constituent Golay sequences; and concatenating at least two of said pairs of constituent Golay sequences to produce the extended sequence.

- 2. A method as claimed in claim 1 wherein the step of transforming the pair of constituent Golay sequences A and B to form each pair of transformed sequences comprises a permutation of elements of the sequences.
- 3. A method as claimed in claim 2 wherein said permutation comprises a reversal of the order of elements in the sequences.
- 4. A method as claimed in claim 2 or 3 wherein said permutation comprises a reversal of a set of address bits identifying locations of elements in the sequences.
- 5. A method as claimed in claim 2 or 3 wherein said permutation comprises a reversal of some but not all of a set of address bits identifying locations of elements in the sequences.
  - 6. A method as claimed in any of claims 2 to 5 wherein said permutation comprises a concatenation of subsets of elements of the sequences selected by decimation.
- 7. A method as claimed in claim 6 wherein there are two subsets of elements, in even and odd locations in the sequences, that are concatenated.
  - 8. A method of producing a preamble for use on an access channel of a cellular communications system, wherein the preamble includes an extended sequence produced by the method of any of claims 1 to 7.
- 25 9. A method as claimed in claim 8 wherein each of the constituent Golay sequences A and B and each of the transformed sequences comprises 256 elements, and the preamble comprises 4096 elements.
- 10. A method as claimed in claim 9 wherein the extended sequence comprises the constituent Golay sequences A and B and seven different transformations of these
   30 sequences A and B.

- 11. A method of providing a preamble for an access channel of a CDMA (code division multiple access) communications system, comprising the steps of: providing a pair of constituent Golay sequences A and B; permuting elements of the pair of sequences A and B to form at least one pair of transformed sequences which are also constituent Golay sequences; and
- concatenating said at least one pair of transformed sequences with the pair of constituent Golay sequences A and B for use as an extended sequence in the preamble.
- 12. A method as claimed in claim 11 wherein the preamble is provided by a concatenation of the pair of constituent Golay sequences A and B with a plurality of pairs of said transformed sequences according to different permutations of the elements of the pair of constituent Golay sequences A and B.
  - 13. A method as claimed in claim 11 or 12 wherein the step of permuting elements of the pair of sequences A and B comprises a reversal of the order of elements in the sequences.
- 15 14. A method as claimed in any of claims 11 to 13 wherein the step of permuting elements of the pair of sequences A and B comprises a reversal of at least some of a set of address bits identifying locations of elements in the sequences.
  - 15. A method as claimed in any of claims 11 to 13 wherein the step of permuting elements of the pair of sequences A and B comprises a concatenation of two subsets of elements of the sequences, the two subsets corresponding respectively to even and odd locations in the sequences.
    - 16. Apparatus for use in carrying out the method of any of claims 1 to 15, comprising a memory for storing elements of each sequence, and address control means for addressing the memory for writing to or reading from the memory consecutively at locations according to the transformed sequences.
  - 17. Apparatus as claimed in claim 16 wherein the address control means comprises a bidirectional counter for supplying memory addresses incremented in either of two opposite directions, and an address permutation unit for permuting said addresses in accordance with the transformed sequences and for supplying the permuted addresses to the memory.

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- 18. A preamble for an access channel of a CDMA (code division multiple access) communications system, comprising a pair of constituent Golay sequences A and B and, concatenated with the pair of constituent Golay sequences A and B, at least one pair of transformed sequences which are also constituent Golay sequences, each pair of transformed sequences comprising a permutation of elements of the pair of sequences A and B.
  - 19. A preamble as claimed in claim 18 comprising a plurality of pairs of said transformed sequences having different respective permutations of the elements of the pair of sequences A and B.
- 10 20. A preamble as claimed in claim 18 or 19 wherein a pair of said transformed sequences comprises the pair of sequences A and B with a reversed order of elements in the sequences.
  - 21. A preamble as claimed in claim 18 or 19 wherein a pair of said transformed sequences comprises the pair of sequences A and B with elements in the sequences permuted in accordance with a reversed order of at least some of a set of address bits identifying locations of elements in the sequences.
    - 22. A preamble as claimed in claim 18 or 19 wherein a pair of said transformed sequences comprises elements in odd locations of the pair of sequences A and B concatenated with elements in even locations in the pair of sequences A and B.

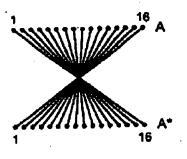


Fig. 1

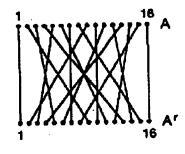


Fig. 2

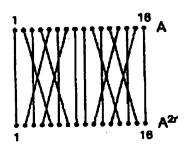


Fig. 3

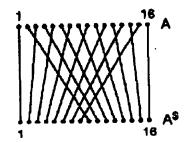


Fig. 4

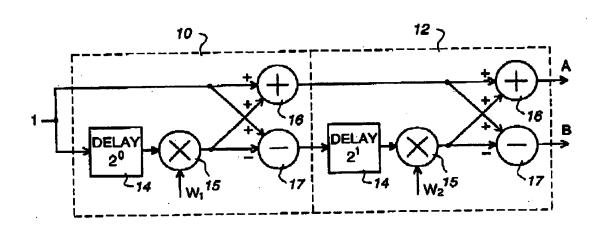
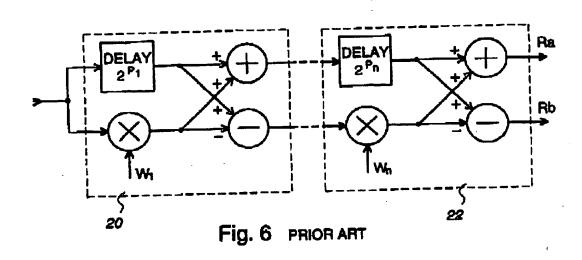


Fig. 5 PRIOR ART



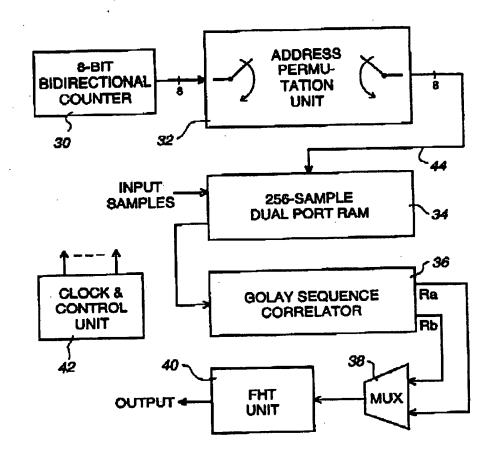


Fig. 7

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